Fig. 1

SEQ ID NO: 1

AGGAAGCCTCAAGACGCGGAGCAGCGCAGGAAGGAGCCCCCGGCAGC CCGGAGGAGCATGGGCACCTTGCGGGATTTACAGTACGCGCTCCAGGA GAAGATCGAGGAGCTGAGCCAGCGGGATGCTCTCATCGACGAGCTGGA GCTGGAGTTGGATCAGAAGGACGAACTGATCCAGAAGCTGCAGAACGA GCTGGACAAGTACCGCTCGGTGATCCGACCAGCCACCCAGCAGCGCCA GAAGCAGAGCGCGAGCACCTTGCAGGGCGAGCCGCGCACCAAGCGGCA GGCGATCTCCGCCGAGCCCACCGCCTTCGACATCCAGGATCTCAGCCA TGTGACCCTGCCCTTCTACCCCAAGAGCCCACAGTCCAAGGATCTTAT AAAGGAAGCTATCCTTGACAATGACTTTATGAAGAACTTGGAGCTGTC GCAGATCCAGGAGATTGTGGATTGTATGTACCCGGTGGAGTATGGCAA GGACAGTTGCATCAAAGAAGAAGGAGACGTGGGGTCACTGGTGTATGT CATGGAAGATGGTAAGGTTGAAGTTACAAAAGAAGGTGTGAAGTTGTG TACCATGGGTCCAGGAAAAGTGTTTGGGGAATTGGCTATTCTTTACAA CTGTACCCGGACAGCGACCGTCAAGACTCTTGTAAATGTAAAACTCTG GGCCATTGATCGACAATGTTTTCAAACAATAATGATGAGGACAGGACT CATCAAGCATACCGAGTATATGGAATTTTTAAAAAGCGTTCCAACATT CCAGAGCCTTCCTGAAGAGATCCTCAGCAAGCTTGCTGATGTCCTTGA AGAGACCCACTATGAAAATGGAGAATATATTATCAGGCAAGGTGCAAG AGGGGACACCTTCTTTATCATCAGCAAAGGAACGGTAAATGTCACTCG TGAAGACTCACCGAGTGAAGACCCAGTCTTTCTTAGAACTTTAGGAAA AGGAGACTGGTTTGGAGAGAAAGCCTTGCAGGGGGAAGATGTGAGAAC *AGCAAACGTAATTGCTGCAGAAGCTGTAACCTGCCTTGTGATTGACAG AGACTCTTTTAAACATTTGATTGGAGGGCTGGATGATGTTTCTAATAA AGCATATGAAGATGCAGAAGCTAAAGCAAAATATGAAGCTGAAGCGGC TTTCTTCGCCAACCTGAAGCTGTCTGATTTCAACATCATTGATACCCT TGGAGTTGGAGGTTTCGGACGAGTAGAACTGGTCCAGTTGAAAAGTGA AGAATCCAAAACGTTTGCAATGAAGATTCTCAAGAAACGTCACATTGT GGACACAAGACAGCAGGAGCACATCCGCTCAGAGAAGCAGATCATGCA

Fig. 1 (continued)

GGGGGCTCATTCCGATTTCATAGTGAGACTGTACAGAACATTTAAGGA CAGCAAATATTTGTATATGTTGATGGAAGCTTGTCTAGGTGGAGAGCT CTGGACCATTCTCAGGGATAGAGGTTCGTTTGAAGATTCTACAACCAG ATTTTACACAGCATGTGTGGTAGAAGCTTTTGCCTATCTGCATTCCAA AGGAATCATTTACAGGGACCTCAAGCCAGAAAATCTCATCCTAGATCA CCGAGGTTATGCCAAACTGGTTGATTTTGGCTTTGCAAAGAAAATAGG ATTTGGAAAGAAACATGGACTTTTTGTGGGACTCCAGAGTATGTAGC CCCAGAGATCATCCTGAACAAAGGCCATGACATTTCAGCCGACTACTG GTCACTGGGAATCCTAATGTATGAACTCCTGACTGGCAGCCCACCTTT CTCAGGCCCAGATCCTATGAAAACCTATAACATCATATTGAGGGGGAT TGACATGATAGAATTTCCAAAGAAGATTGCCAAAAATGCTGCTAATTT AATTAAAAAACTATGCAGGGACAATCCATCAGAAAGATTAGGGAATTT GAAAAATGGAGTAAAAGACATTCAAAAGCACAAATGGTTTGAGGGCTT TAACTGGGAAGGCTTAAGAAAAGGTACCTTGACACCTCCTATAATACC AAGTGTTGCATCACCCACAGACACAAGTAATTTTGACAGTTTCCCTGA GGACAACGATGAACCACCACCTGATGACAACTCAGGATGGGATATAGA CTTCTAATGTATTTCTCTTACCTGCTTCTGCCTTGCTGAAGACAGCTT GTGCTCGGGGTCACCATGATGCCTTTGATCGATGCTGCTCCAGTAACT ACAGTGGCATTAGGACTTATCGCTTAGATGACAATAGTGCTCTTTACA TGTTTTCTGTTTGAACCTAAAATAGCAGTTGACATGGTGGTCCTGAAG CAAAGCCTTTCACCAGTAAAGAGATGTTTTCTATTGTTGCAATGACCT TGCTTTGCTCTGATTATAATTTGAAAGACTGTAGGAAACACTTCAATG TAGTATAAGAGTCTGTACCTTGCTGGAATATTCAAGAAGATGAAAGAA TAATATTGGGTACAATAGATTACTATGGTACAGAAACTGGGCTATT CCCTTTCTTCAAGTGAAGGCTGTGGGATCTATTACAGCTGCAGGCCGG TGTATATACCATACAAAAGAGGACCACACATCTGTTGGTCACAGAGTT CATGTCACACCAGTGCTAGAAGTTTCATGATTTTATTTCCCAGCAGTG CTGATGACAAGACTGAATGTTACCTTTTCTTTCTGACAGATTTTAAAA

Fig. 1 (continued)

ATTGATATGATAAAAGCACAACTGCTATAGATŢCTGCTGAGACCTCTC ATAGTAGGTATATATGAGTTTTCACAGAAGACTGAAAAATAATGCATG ATATTTGTTTTTTTTTTGATAAATTGGCATGACAGAGTGGGGAAAA AAAGCAATTCACAAAACCATTTCATATTTTTTTAAAATATTGTGCTTAA AGATGGTCCTGGAAGTAAATGACTAGCAGCCAATTGGTTTTACTTAAC CTTGGGGTAGGGT_GGAGTGGGGAGGGATTAAAACCCATCCAAAAAATA AATAAAAACTATATAGGTGCTATGTATATCTTTCATCTGTAAATGTCA GTGTCTGAACAGCAACACAAATTCAAATCATTATACGTGTAGCCAGAA ACTCAAGCATTTTCACTAAAGTTATTAAACCAAACTCCTGTCCAATTT GACTTATACAACATAGTCAGTCTAGAGTTGAGAGACAAAGGTAATTAT AAACCTATTTGAACTAGCTTCTTGTCTTAGGCCTGAACCAAAAAACAA CAAACAAACAAAAAACAAGAATGAAAAACAGAAATAAAAGAAGTAGAA AAGACAAAGAAAGCCCCAAAGTCAAAGTTGTTAATATTTTACAGGT TTACCAGATCTGGAACATTACTTATTTGAGGTCAGAGAACAAAACAAG AACCTGGCCAGGTGTTGATTACCTTTTAGTGAATAAGCTGAGTCCATA TACTTGTCTAACTAAGAAAGCAGTACAGAGGAAAACAGGAACCTGATT TTTTTAAAATAAATTTTAAATAAAATAGAATTACTACAATTCTGCAAT TTCATACTACCTAAAAAAGACTAGATTTGAAAATGTCAAGCTGATTTA CTTTATTCACATGGAGAAAAGAATCCACAAATTAAACTGAGTCCTTCA CTGGCATGCCAGTTGACTATTATTAGCTGTCATAAGTAACCCCG

Fig. 2

SEQ ID NO: 2

MGTLRDLQYALQEKIEELRQRDALIDELELELDQKDELIQKLQNELDK
YRSVIRPATQQAQKQSASTLQGEPRTKRQAISAEPTAFDIQDLSHVTL
PFYPKSPQSKDLIKEAILDNDFMKNLELSQIQEIVDCMYPVEYGKDSC
IIKEGDVGSLVYVMEDGKVEVTKEGVKLCTMGPGKVFGELAILYNCTR
TATVKTLVNVKLWAIDRQCFQTIMMRTGLIKHTEYMEFLKSVPTFQSL
PEEILSKLADVLEETHYENGEYIIRQGARGDTFFIISKGTVNVTREDS
PSEDPVFLRTLGKGDWFGEKALQGEDVRTANVIAAEAVTCLVIDRDSF
KHLIGGLDDVSNKAYEDAEAKAKYEAEAAFFANLKLSDFNIIDTLGVG
GFGRVELVQLKSEESKTFAMKILKKRHIVDTRQQEHIRSEKQIMQGAH
SDFIVRLYRTFKDSKYLYMLMEACLGGELWTILRDRGSFEDSTTRFYT
ACVVEAFAYLHSKGIIYRDLKPENLILDHRGYAKLVDFGFAKKIGFGK
KTWTFCGTPEYVAPEIILNKGHDISADYWSLGILMYELLTGSPPFSGP
DPMKTYNIILRGIDMIEFPKKIAKNAANLIKKLCRDNPSERLGNLKNG
VKDIQKHKWFEGFNWEGLRKGTLTPPIIPSVASPTDTSNFDSFPEDND
EPPPDDNSGWDIDF

<u>Fig. 3</u>

SEQ ID NO: 3

5'- agccgactactggtcactgg -3'

Fig. 4

SEQ ID NO: 4

5'- gatctgggcctgagaaaggt -3'

Fig. 5

SEQ ID NO: 5

5'- tgtatgaactcctgactggcagccc -3'